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	L3	satellite tobacco necrosis virus	31
	L2	satellite tobacco mosaic virus	11
	DB=US	PT; PLUR=YES; OP=ADJ	
	L1	satellite tobacco mosaic virus	5

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                 alerts (SDIs) affected
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      12 DEC 17
                 CERAB reloaded; updating to resume; current-awareness
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NEWS
                 February 2005
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      17 FEB 25
                 CA/CAPLUS - Russian Agency for Patents and Trademarks
                  (ROSPATENT) added to list of core patent offices covered
      18 FEB 10
                 STN Patent Forums to be held in March 2005
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NEWS
      19 FEB 16
                 STN User Update to be held in conjunction with the 229th ACS
                 National Meeting on March 13, 2005
NEWS 20 FEB 28 PATDPAFULL - New display fields provide for legal status
                 data from INPADOC
NEWS 21 FEB 28 BABS - Current-awareness alerts (SDIs) available
NEWS 22 FEB 28 MEDLINE/LMEDLINE reloaded
NEWS 23 MAR 02 GBFULL: New full-text patent database on STN
NEWS 24 MAR 03 REGISTRY/ZREGISTRY - Sequence annotations enhanced
NEWS 25 MAR 03 MEDLINE file segment of TOXCENTER reloaded
NEWS EXPRESS JANUARY 10 CURRENT WINDOWS VERSION IS V7.01a, CURRENT
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=> file agricola caplus biosis

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=> s satellite tobacco mosaic virus L1 137 SATELLITE TOBACCO MOSAIC VIRUS

=> s l1 and coat protein

L3 37 L1 AND COAT PROTEIN

=> dup rem 13

PROCESSING COMPLETED FOR L3

L4 18 DUP REM L3 (19 DUPLICATES REMOVED)

=> d 1-10 ti

- L4 ANSWER 1 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- TI Molecular structures of viruses from Raman optical activity.
- L4 ANSWER 2 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN Stability of Satellite Tobacco Mosaic Virus RNA cores and its implications for infectivity and macromolecular assembly.
- L4 ANSWER 3 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 2
- TI Biophysical studies on the RNA cores of satellite tobacco mosaic virus
- L4 ANSWER 4 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 3
- TI Heterogeneity in the 3'-terminal untranslated region of tobacco mild green mosaic tobamoviruses from Nicotiana glauca resulting in variants with three or six pseudoknots

- L4 ANSWER 5 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 4
- TI Naturally occurring variants of satellite tobacco mosaic virus.
- L4 ANSWER 6 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 5
- TI Refined structure of satellite tobacco mosaic virus at 1.8 Å resolution
- L4 ANSWER 7 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 6
- TI Replication of wild-type and mutant clones of **satellite tobacco mosaic virus** in Nicotiana benthamiana protoplasts.
- L4 ANSWER 8 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN Efficient replication of mutants of satellite tobacco mosaic virus in Nicotiana protoplasts.
- L4 ANSWER 9 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN TI The structure of satellite panicum mosaic virus at 1.9 A resolution.
- L4 ANSWER 10 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 7
- TI Structural comparison of the plant satellite viruses.
- => d 2 ab
- L4 · ANSWER 2 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- => d 2 so
- L4 ANSWER 2 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN SO Biophysical Journal, (January, 2002) Vol. 82, No. 1 Part 2, pp. 461a. print.

Meeting Info.: 46th Annual Meeting of the Biophysical Society. San Francisco, California, USA. February 23-27, 2002. CODEN: BIOJAU. ISSN: 0006-3495.

- => d 7 ab
- L4 ANSWER 7 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 6
- AB RNA transcribed from cloned satellite tobacco
  mosaic virus (STMV) cDNA replicated in Nicotiana
  benthamiana protoplasts when co-inoculated with tobacco mild green mosaic
  virus (TMGMV) genomic RNA, but degraded when inoculated alone. STMV
  genomic RNA extracted from wild-type virions replicated in protoplasts
  when co-inoculated with TMGMV, tobacco mosaic virus (TMV) or tomato mosaic
  virus (ToMV). Transcripts from clones of two STMV coat
  protein (CP) mutants accumulated to the same level as wild-type
  transcripts in protoplasts when co-inoculated with TMGMV, whereas a third
  mutant accumulated to detectable levels in some, but not all, experiments.
  These results confirm that STMV RNA requires helper virus for replication,
  and that the helper specificity exhibited by cloned STMV reflects a
  specific requirement for the TMGMV replicase. It also demonstrates that

the low accumulation of STMV CP mutants observed previously in whole plants cannot be attributed to inefficient RNA replication.

#### => d 11-18 ti

- ANSWER 11 OF 18 AGRICOLA Compiled and distributed by the National L4Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. **DUPLICATE 8** (2005) on STN
- Characterization of deletion and frameshift mutants of satellite TItobacco mosaic virus
- **L4** ANSWER 12 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2005) on STN DUPLICATE 9
- Similarities between the secondary structure of satellite ТT tobacco mosaic virus and tobamovirus RNAs.
- ANSWER 13 OF 18 AGRICOLA Compiled and distributed by the National L4Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2005) on STN DUPLICATE 10
- ΤI Three-dimensional structure of satellite tobacco mosaic virus at 2.9 angstroms resolution.
- ANSWER 14 OF 18 AGRICOLA Compiled and distributed by the National L4Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2005) on STN DUPLICATE 11
- ТT Double-helical RNA in satellite tobacco mosaic virus.
- ANSWER 15 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN L4
- ТT Recombinant expression system based on satellite tobacco mosaic virus
- ANSWER 16 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on L4
- TΙ THE COMPLETE NUCLEOTIDE SEQUENCE OF THE GENOMIC RNA OF THE TOBAMOVIRUS TOBACCO MILD GREEN MOSAIC VIRUS.
- ANSWER 17 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN L4
- TI Nucleotide sequence and translation of satellite tobacco mosaic virus RNA
- ANSWER 18 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 12 L4
- ΤI Analysis of the genome of satellite panicum mosaic virus

# => d 11 ab

- ANSWER 11 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2005) on STN DUPLICATE 8
- AR A series of frameshift and deletion mutations was created in the genome of satellite tobacco mosaic virus (STMV) by modifying full-length cDNA clones of the type strain, from which biologically active transcripts could be synthesized in vitro. Deletions and frameshift mutations in the 5' open reading frame had no effect compared to wild-type STMV, on RNA accumulation, systemic movement or the symptoms induced by STMV in Nicotiana tabacum co-inoculated with tobacco mild green mosaic tobamovirus (TMGMV). This implies that the protein encoded by this reading frame is not necessary for biological activity. Deletions and frameshift mutations in the coat protein

open reading frame resulted in decreased accumulation of STMV RNA in N.

tabacum, although these mutants were still capable of systemic movement, presumably in a nonencapsidated or free RNA form. Furthermore, the mild symptoms induced in tobacco by co-inoculations of wild-type STMV/TMGMV or infection with TMGMV alone were altered to severe systemic necrosis when plants were co-inoculated with these STMV coat protein mutants and TMGMV. Mutants within the 3' untranslated region were much less able to accumulate in TMGMV-infected plants than was wild-type STMV, and under some growth conditions did not accumulate to detectable levels.

# => d 12 ab

- L4 ANSWER 12 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 9
- AB The secondary structure of satellite tobacco
  mosaic virus (STMV) RNA was predicted using computer
  simulations of RNA folding. The analogies of structural elements in the 3'
  end untranslated regions (3'-UTR) of tobamoviral RNAs were analysed. In
  addition to the tRNA-like structure and pseudoknot stalk, which are found
  in all known RNAs of tobamoviruses and STMV, another region of stable
  consecutive pseudoknots was predicted in the 3'-UTR of STMV RNA. A similar
  pattern of repeated structural units, containing pseudoknot stalks and
  parts of the tRNA-like structure, was also found in odontoglossum ringspot
  virus (ORSV) RNA 3'-UTR. The predictions on the structure are supported by
  sequence comparisons which point to an important functional role of 3'
  terminal pseudoknots in STMV RNA as well as in other tobamoviral RNAs. The
  possible participation of pseudoknotted structures in the interactions
  with coat protein in STMV is discussed.

#### => d 15 ab

- L4 ANSWER 15 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
- AB A cDNA encoding a full-length copy of the RNA genome of satellite tobacco mosaic virus is cloned and characterized for use as a transformation vector for plants. The virus is systemic and needs a helper virus for replication and so can be used to introduce foreign DNA into a plant without being pathogenic. The genome was cloned as a cDNA by standard methods using defined oligonucleotide primers. The RNA was shown by in vitro translation to encode two peptides of mol. weight 6,800 and 17,500. A series of plasmid derivs. containing modified cDNAs were constructed and transcripts from these used in infectivity studies on tobacco plants with tobacco mosaic virus isolates as helpers. The transcripts tested produced infected plants and virions were recovered from these plants. Antisense transcripts were not infective.

## => d 15 so

L4 ANSWER 15 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN SO PCT Int. Appl., 48 pp.

CODEN: PIXXD2

=> d 15 pi

L4 ANSWER 15 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
PATENT NO. KIND DATE APPLICATION NO. DATE

PI WO 9012107 A1 19901018 WO 1990-US1738 19900402
W: CA, JP, US
RW: AT, BE, CH, DE, DK, ES, FR, GB, IT, LU, NL, SE

=> d 17 ab

AΒ

L4 ANSWER 17 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN

Satellite tobacco mosaic virus (STMV) is a plant virus with a 17-nm icosahedral particle encapsidating a 0.3 + 106 Mr ssRNA genome that depends on tobamoviruses for its replication. The complete nucleotide sequence of STMV RNA deduced in the expts. described here was 1059 nucleotides in length. The efficiency of labeling viral RNA with  $[\gamma-32P]$ ATP using T4 polynucleotide kinase was not affected by treatment with tobacco acid pyrophosphatase and/or bacterial alkaline phosphatase, indicating that the majority of the 5' termini of encapsidated STMV RNAs were phosphorylated. The 240 3'terminal nucleotides of STMV RNA and either tobacco mosaic virus (TMV) Ul RNA or TMV U2/U5 RNA had greater than 65% overall sequence similarity, with 2 nearly identical regions of 40 and 50 bases, resp. There were no other regions of sequence homol. with TMV RNA. The 19 5'-terminal nucleotides of STMV RNA had greater than 65% sequence similarity with the 16 5'-terminal nucleotides of brome mosaic virus RNA 3 and 50% sequence similarity with the 12 5'-terminal nucleotides of the Q strain of cucumber mosaic virus RNA 3. The first open reading frame (ORF) beginning at base 53 encoded a 6800 Mr protein that corresponded in size to a major in vitro translation product directed by STMV RNA. A second ORF, beginning at nucleotide 163, had the capacity to code for a protein that corresponded in size (17,500 Mr) to the other major in vitro translation product. The first 12 codons of this ORF corresponded to the sequence of the N-terminal amino acids of the capsid protein. Western blot anal. of the in vitro translation products revealed that the 17,500 Mr protein had the same electrophoretic mobility as the authentic capsid protein; it was also antiquenically related to the capsid protein, but the 6800 Mr protein was not. Time course anal. of in vitro translation demonstrated that the 6800 Mr protein was synthesized at the same time as the capsid protein and did not arise by the proteolytic cleavage of a larger precursor polypeptide. These results suggest that the genome of STMV functioned as a polycistronic mRNA. It has not been determined if the 6800 Mr protein is synthesized in vivo. STMV RNA had untranslated regions of 52 and 418 nucleotides at its 5' and 3' termini, resp. Nonphosphorylated 5' termini, the degree of similarity to the 3' terminus of 2 of its helper viruses, the genome organization, and the ability to function as a polycistronic mRNA are unique features for the genome of this satellite virus.

=> d 17 so

L4 ANSWER 17 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN SO Virology (1989), 170(1), 139-46 CODEN: VIRLAX; ISSN: 0042-6822

=> s satellite tobacco necrosis virus
L5 316 SATELLITE TOBACCO NECROSIS VIRUS

=> s 15 and vector
L6 14 L5 AND VECTOR

=> dup rem 16
PROCESSING COMPLETED FOR L6
L7 9 DUP REM L6 (5 DUPLICATES REMOVED)

=> d 1-9 ti

L7 ANSWER 1 OF 9 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN Geminate structures of African cassava mosaic virus.

L7 ANSWER 2 OF 9 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN

- TI 1966-1996: Thirty years of virology.
- L7 ANSWER 3 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- TI Expression of tobacco necrosis virus open reading frames 1 and 2 is sufficient for the replication of satellite necrosis virus.
- L7 ANSWER 4 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 2
- TI Evidence that the requirements for ATP and wheat germ initiation factors 4A and 4F are affected by a region of **satellite tobacco necrosis virus** RNA that is 3' to the ribosomal binding site
- L7 ANSWER 5 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 3
- TI Satellite tobacco necrosis virus:
  a new vector in plant genetic engineering.
- L7 ANSWER 6 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Synthesis of satellite tobacco necrosis
  virus (STNV) RNA-like transcripts in Escherichia coli
- L7 ANSWER 7 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 4
- TI The determination of the heavy-atom substitution sites in the satellite tobacco necrosis virus
- L7 ANSWER 8 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Progress toward a low resolution structure of the satellite tobacco necrosis virus
- L7 ANSWER 9 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Amino acid composition, antigenicity, and other characterisits of the satellite viruses of tobacco necrosis virus

## => d 3 ab

- ANSWER 3 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- AB Tobacco necrosis virus (TNV) is a small icosahedral plant virus which is often associated with satellite viruses. The genomic RNA of TNV contains six open reading frames (ORFs), of which ORFs 1 and 2 are thought to encode the viral polymerase. We demonstrate that tobacco protoplasts transfected with a **vector** containing TNV ORFs 1 and 2 under the control of the cauliflower mosaic virus 35S promoter, as well as protoplasts derived from transgenic Nicotiana tabacum containing the same gene(s), support replication of **satellite tobacco**necrosis virus RNA.

# => d 3 so

- L7 ANSWER 3 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- SO Virology, Sept 10, 1995. Vol. 212, No. 1. p. 222-224 Publisher: Orlando, Fla. : Academic Press. CODEN: VIRLAX; ISSN: 0042-6822

L7 ANSWER 5 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

(2005) on STN DUPLICATE 3

=> d 5 so

- L7 ANSWER 5 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 3
- SO Symbiosis, 1986. Vol. 2, No. 1/3. p. 35-41 Publisher: Philadelphia : Balaban Publishers. ISSN: 0334-5114

## => 6 ab

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ANSWER 6 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN

=> d 6 ab

L7

- AB By means of genetic engineering, a full-size double-stranded DNA copy of the STNV RNA genome was inserted into a prokaryotic expression plasmid, so that transcription of the STNV information, starting at the PL-promoter, can be induced. Although no Shine-Dalgarno region (bacterial ribosome-binding site) is present in the original expression vector, induction of transcription resulted in the synthesis of STNV coat-protein. By analyzing the 5'-untranslated region of the STNV genome, a shine-Dalgarno-like sequence could be found. Anal. of the
  - genome, a shine-Dalgarno-like sequence could be found. Anal. of the transcripts by RNA extraction, followed by Northern blotting and hybridization with radiolabeled STNV-DNA fragments revealed that a great deal of the transcripts terminate at or close to the 3' end of the STNV information. Whether this phenomenon represents a transcription stop or a modification site still remains to be solved.

=> dup rem 18
PROCESSING COMPLETED FOR L8
L9 26 DUP REM L8 (20 DUPLICATES REMOVED)

=> d 1-10 ti

- L9 ANSWER 1 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Role of viral movement and coat proteins and RNA in phloem-dependent movement and phloem unloading of tobamoviruses
- L9 ANSWER 2 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Identification of gene function in plants by gene silencing and the use of viral satellites to deliver silencing RNA
- L9 ANSWER 3 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Replication-incompetent virus-like particles for targetted delivery of nucleic acids or proteins
- L9 ANSWER 4 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- TI A conserved, precise RNA encapsidation pattern in tobamovirus particles.

- L9 ANSWER 5 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Packaging of RNA into viral particles
- L9 ANSWER 6 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 2
- TI Expression of tobacco mosaic virus coat protein and assembly of pseudovirus particles in Escherichia coli.
- L9 ANSWER 7 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 3
- TI Uncoating of tobacco mosaic virus RNA in protoplasts.
- L9 ANSWER 8 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Minimum sequence requirement for encapsidation of the cowpea strain of tobacco mosaic virus
- L9 ANSWER 9 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 4
- TI Mutation and replacement of the 16-kDa protein gene in RNA-1 of tobacco rattle virus.
- L9 ANSWER 10 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 5
- TI Study of TMV assembly with heterologous RNA containing the **origin** -of-assembly sequence

# => d 2 pi

L9	ANSWER PATENT												NO.		Dž	ATE	
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ΡI	WO 2000063397			•	A2 20001026 WO 2000-EP3521							20000417					
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# => d 4 gb

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L9 ANSWER 4 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

(2005) on STN DUPLICATE 1

The bidirectional RNA encapsidation pathway in nine sequenced Type 1 AB Tobamovirus genomes will result in RNA-coat protein assembly, up to and including the first transcribed G, adjacent to the 5'-cap structure (m7 Gppp). This precision is highly conserved, despite wide interstrain variations in the absolute position of the phase-determining core of the origin-of-assembly sequence (Gxx)n and in overall genome length (6311-6507 nts). A Type 2 Tobamovirus genome did not comply with this pattern. All genomes had a statistically significant bias for G at every third (or 3n) position, resulting in a preponderance of GNN codons and hence a high Val, Ala, Gly, Asp, Glu content, at least in the large (126/183 kDa) and amino-coterminal replicase protein genes. Contrary to predictions from the X-ray fibre diffraction structure of tobacco mosaic virus (TMV, U1 strain), only one (pepper mild mottle virus) of the nine Type 1 Tobamoviruses positioned the preferred G-repeat in the most favourable (5') position of the trinucleotide binding site on each coat protein (CP) subunit. In all but one of the eight remaining Type 1 Tobamovirus genomes, G would predominate in the CP 3'-site. The significance of these observations for TMV particle assembly, disassembly and host cell interactions are discussed.

## => d 4 so

- L9 ANSWER 4 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- SO Archives of virology, 1995. Vol. 140, No. 9. p. 1677-1685 Publisher: Wien, Austria : Springer-Verlag. CODEN: ARVIDF; ISSN: 0304-8608

# => d 5 ab

L9 ANSWER 5 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN

An in vivo system for expression and packaging of RNA into pseudovirus ΑB particles is described. The invention is based on the discovery that genes for plant viral coat proteins (CPs) may be efficiently expressed in E. coli, and that these coat proteins will assemble in vivo and package chimeric RNA, containing an origin-of-assembly (OAS) sequence, to form mature viral particles containing a foreign RNA. present invention thus provides for packaging of RNA into a RNase-resistant form that is easily purified and stored, and which overcomes the prior art problems associated with the degradation of RNA by RNases. Significantly, the method of the invention is RNA sequence- and length-independent. The components of the invention include a source of viral coat proteins, and an expression cassette directing transcription of DNA encoding an OAS-containing transcript. The CPs and OAS are from a plant virus having a rod-shaped helical particle and a single-stranded RNA genome, most preferably tobacco mosaic virus

# => d 5 pi

L9	ANSWER	5 OF	26	CAP	LUS	COP	YRIG	HT 2	005	ACS	on S'	TN					
	PATENT	NO.			KIN	)	DATE			APPL	ICAT	ION I	. O <i>l</i>		D	ATE	
						-									-		
PΙ	WO 9410329				A1	A1 19940511				WO 1993-US10396					19931028		
•	W:	AU,	BB,	BG,	BR,	BY,	CA,	CZ,	FI,	HU,	JP,	KR,	ΚZ,	LK,	LV,	MG,	MN,
		MW,	NO,	NZ,	PL,	RO,	RU,	SD,	SK,	UA,	UΖ,	VN					
	RW:	AT,	ΒĒ,	CH,	DE,	DK,	ES,	FR,	GB,	GR,	ΙE,	IT,	LU,	MC,	NL,	PT,	SE,
		BF,	ВJ,	CF,	CG,	CI,	CM,	GA,	GN,	ML,	MR,	NE,	SN,	TD,	TG		
	US 5443	969			Α		1995	0822		US 1	992-	9711	01		1	9921	029
	AU 9454	554			A1		1994	0524		AU 1	994-	5455	4		1	9931	028
	AU 6937	70			B2		1998	0709									
	EP 6838	21			A1		1995	1129		EP 1	993-	9251	19		1	9931	028

R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LI, LU, MC, NL, PT, SE JP 08505523 T2 19960618 JP 1993-511342 19931028

=> d 8 ab

L9 ANSWER 8 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN

AB Unavailable

=> d 8 so

L9 ANSWER 8 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN

SO (1992) 121 pp. Avail.: Univ. Microfilms Int., Order No. DA9310697 From: Diss. Abstr. Int. B 1993, 53(12, Pt. 1), 6138

=> d 10 ab

AB

L9 ANSWER 10 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 5

The assembly of tobacco mosaic virus (TMV) in initiated by a specific reaction between a capsid protein oligomer and an origin-of-assembly region (OAS) located 900 nucleotides from the 3' terminus of virion RNA. Packaging is then completed by rod elongation both in the 5' and 3' directions. The temporal order of the direction of elongation and the characteristics of the reaction were studied by anal. of the in vitro assembly reaction between strain U1 protein oligomers and transcripts containing a strain U1 OAS embedded at different positions in heterologous RNA. The results confirm that elongation in the 5' direction starts very soon after the initiation reaction and is completed rapidly, within minutes. Packaging in the 3' direction is slower and does not appear to commence until 5' rod formation is complete. The reaction of strain U2 protein with the strain U1 OAS initiates rapidly, but elongation occurs only in the 5' direction; 3' packaging does not occur except when the OAS is at or near the 5' terminus, in which case elongation in the 3' direction initiates without delay with either the U1 or U2 protein. Pauses occur during elongation in the 3' direction at an average of 320 nucleotides, indicating a packaging periodicity of about six to eight helical turns.

=> d 10 so

L9 ANSWER 10 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 5

SO Virology (1990), 174(2), 337-44 CODEN: VIRLAX; ISSN: 0042-6822

=> d 11-20 ti

- L9 ANSWER 11 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Direct recovery of in vitro transcripts in a protected form suitable for prolonged storage and shipment at ambient temperatures
- L9 ANSWER 12 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 6
- TI Assembly of hybrid RNAs with tobacco mosaic virus coat protein. Evidence for incorporation of disks in 5'-elongation along the major RNA tail
- L9 ANSWER 13 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 7
- TI Tobacco mosaic virus coat protein and reporter gene transcripts containing the TMV origin-of-assembly sequence do not interact in double-transgenic tobacco plants: implications for coat protein-medicated protection.

- L9 ANSWER 14 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Hybrid RNA virus which allows encapsidation of recombinant viral sequences in heterologous protein capsids, and their use in plant genetic engineering
- L9 ANSWER 15 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 8
- TI Selective recovery of foreign gene transcripts as virus-like particles in TMV-infected transgenic tobaccos
- L9 ANSWER 16 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 9
- TI Selective encapsidation of CAT gene transcripts in TMV-infected transgenic tobacco inhibits CAT synthesis
- L9 ANSWER 17 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Recombinant RNA packaging system using tobacco mosaic virus
- L9 ANSWER 18 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 10
- TI The effect of multiple dispersed copies of the **origin**-of **assembly** sequence from TMV RNA on the morphology of pseedovirus particles assembled in vitro
- L9 ANSWER 19 OF 26 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 11
- TI Packaging of recombinant RNA molecules into pseudovirus particles directed by the origin-of-assembly sequence from tobacco mosaic virus RNA.
- L9 ANSWER 20 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 12
- TI Addressed fragmentation of tobacco mosaic virus RNA: excision of the 3'-proximal region containing the coat protein gene

## => d 12 ab

singly or a few at a time.

- L9 ANSWER 12 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 6
- During the reassembly of tobacco mosaic virus AB (TMV) RNA, with the coat protein supplied as a disk preparation, the lengths of RNA protected from nuclease are quantized with steps which correspond to incorporation of the subunits from either a single or, more commonly, both rings of a disk. This interpretation has been challenged and it was suggested that the pattern was due to special, though unspecified features of the sequence of TMV RNA. To test whether the specific sequence of TMV RNA is important during the elongation, rather than just during nucleation, growth of particles containing hybrid RNAs was followed with the TMV RNA origin of assembly but otherwise non-TMV sequences. RNA transcripts containing heterologous RNA 5' to the origin of assembly sequence from TMV RNA, i.e. with a heterologous RNA tail in place of the natural major 5'-tail and no minor tail, were prepared in vitro and used for assembly expts. In each case a banding pattern was found that was very similar to that found with native TMV RNA and with a dominant quantum step of just over 100 bases, and sometimes also a step of 50 bases, strongly suggesting that this is not due to any feature of the TMV RNA. This same repeat is also visible even with a heterologous RNA chosen because it had a sequence repeat of 135 or 136 bases, confirming that the quantization is due to a feature of the elongation reaction and in no way to the RNA sequence being encapsidated. Elongation was also followed with the origin of assembly located 5' to the heterologous RNA. This leads to a slower elongation along this 3'-tail, after the initial rapid encapsidation of the origin RNA, which lacks any quantization of length protected. These results are fully compatible with the hypothesis advanced earlier, that the major growth along the 5'-tail is from preformed aggregates (disks) while the minor growth along the 3'-tail is from subunits in the A-protein adding

#### => d 12 so

- L9 ANSWER 12 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 6
- SO Journal of Molecular Biology (1989), 209(3), 407-22

CODEN: JMOBAK; ISSN: 0022-2836

## => d 14 b

'B' IS NOT A VALID FORMAT

In a multifile environment, a format can only be used if it is valid in at least one of the files. Refer to file specific help messages or the STNGUIDE file for information on formats available in individual files.

REENTER DISPLAY FORMAT FOR ALL FILES (FILEDEFAULT): ab

- L9 ANSWER 14 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- AB A hybrid viral RNA comprising an infectious viral sequence, a heterologous origin of assembly and coat protein gene (e.g. derived from a rod-shaped virion), and a heterologous sequence encoding a functional protein can be used to transfect a plant and prepare genetically altered plants. Plasmid pB3RS2, containing a full-length cDNA copy of the brome mosaic virus (BMV) RNA3 fused to the phage T7 promoter, was prepared A 0.5 kb fragment in the middle of coat protein cDNA was removed and replaced with a 0.6 kb fragment containing coat protein-encoding cDNA of tobacco mosaic virus (TMV). This sequence also included the encapsidation origin of the RNA. Barley protoplasts were inoculated with transcripts of this linearized plasmid and of BMV RNA1 and RNA2 cDNAs. The infected cells produced TMV virion particles which looked normal (serol. specific electron microscopy).

# => d 14 so

- L9 ANSWER 14 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- SO Eur. Pat. Appl., 9 pp.

CODEN: EPXXDW

#### => d 14 pi

L9	ANSWER 14 OF 26 PATENT NO.	CAPLUS (	COPYRIGHT :	2005 ACS on STN APPLICATION NO.	DATE
ΡI	EP 278667	A2	19880817	EP 1988-300908	19880203
	EP 278667	A3	19900905		
	EP 278667	B1	19940720		
	R: AT, BE,	CH, DE, ES	FR, GB,	GR, IT, LI, LU, NL, S	SE
	ES 2060646	Т3	19941201	ES 1988-300908	19880203
	AU 8811383	A1	19880811	AU 1988-11383	19880208
	AU 606382	B2	19910207		
	CA 1337933	A1	19960116	CA 1988-558357	19880208
	JP 63301787	A2	19881208	JP 1988-28630	19880209
	JP 3479531	B2	20031215		
	US 5602242	Α	19970211	US 1995-445990	19950522
	US 5627060	Α	19970506	US 1995-473617	19950607
	US 5804439	Α	19980908	US 1996-773821	19961227
	JP 2004000260	A2	20040108	JP 2003-276352	20030717

### => d 17 ab

- L9 ANSWER 17 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- AB A chimeric RNA was produced which contained the origin of assembly sequence (responsible for packaging) of tobacco mosaic virus (TMV) and a sequence coding for a foreign protein. The chimeric RNA was made by producing a cDNA copy of the

origin of assembly, a cDNA for the foreign protein, ligating these 2 together in a suitable plasmid, and transcribing them in vitro. The transcripts were packaged into pseudoviral particles in an in vitro system that had a ratio of TMV coat protein to RNA of 100:1. These pseudoviruses can infect a wide variety of plant systems and even Escherichia coli and Xenopus laevis cells, expressing the foreign protein in these systems. Sequences for foreign proteins that were incorporated into the recombinant RNA packaging system were those coding for calf prepropchymosin, chicken lysozyme, X. borealis rRNA, and transposon Tn9 chloramphenicol acetyltransferase.

=> d 17 so

L9 ANSWER 17 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN

SO PCT Int. Appl., 36 pp.

CODEN: PIXXD2

=> d 17 pi

L9 ANSWER 17 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
PATENT NO. KIND DATE APPLICATION NO. DATE
PI WO 8706261 A1 19871022 WO 1987-GB249 19870413
W: GB, JP, US

RW: AT, BE, CH, DE, FR, GB, IT, LU, NL, SE

=> d 19 ab

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(2005) on STN DUPLICATE 11

=> d 19 so

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  (2005) on STN DUPLICATE 11
- SO Virology, Dec 1986. Vol. 155, No. 2. p. 299-308 ill Publisher: Duluth, Minn. : Academic Press. CODEN: VIRLAX; ISSN: 0042-6822

#### => d 21-26 tui

'TUI' IS NOT A VALID FORMAT

In a multifile environment, a format can only be used if it is valid in at least one of the files. Refer to file specific help messages or the STNGUIDE file for information on formats available in individual files.

REENTER DISPLAY FORMAT FOR ALL FILES (FILEDEFAULT):ti

- L9 ANSWER 21 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 13
- TI Tobacco mosaic virus induces the synthesis of a family of 3'-coterminal messenger RNAs and their complements
- L9 ANSWER 22 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 14
- TI A study of TMV ts mutant Ni2519. III. Location of the reconstitution initiation sites on Ni2519 RNA
- L9 ANSWER 23 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Location of reconstitution initiation sites of the tobacco mosaic virus mutant Ni2519 RNA

- L9 ANSWER 24 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 15
- TI The nucleotide sequence at the origin for assembly on tobacco mosaic virus RNA
- L9 ANSWER 25 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Location of the origin for viral reassembly on tobacco mosaic virus RNA and its relation to stable fragment
- L9 ANSWER 26 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN
- TI **Tobacco mosaic virus** assembly specificity and the transition in protein structure during RNA packaging

=> d 24 ab

AB

L9 ANSWER 24 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 15

RNA extracted from tobacco mosaic virus (TMV) assembly nucleation complexes consists of a population of fragments 50-550 nucleotides in length. The shortest fragments define a core .apprx.100 residues long common to all the fragments, whereas larger ones are covalently extended by ≤400 nucleotides in 1 direction and  $\leq$ 30 in the other. The most probable interpretation is that assembly nucleates at a unique internal site on the RNA, with growth following bidirectionally but at greatly unequal rates. A nucleotide sequence of 149 residues around the assembly origin was determined The sequencing data are consistent only with a 3' to 5' major direction of growth. Unusual features occur both within the sequence of the 1st disk binding site and elsewhere in the nucleation region; these may be involved in directing the binding and(or) dislocation of ≤3 coat protein disks. The nucleotide sequence around the assembly origin is homologous to, but distinct from, part of the nucleotide sequence from the TMV coat protein gene. The assembly origin may itself have a dual role as a coding region within another gene. There is striking clustering of nucleotide residues that show homol. between the assembly origin and the coat protein gene in those regions coding (in the coat gene) for conserved amino acids, although the protein sequences that may be translated from the 2 genes are dissimilar.

=> d 24 so

=> dis his

L9 ANSWER 24 OF 26 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 15 SO Cell (Cambridge, MA, United States) (1977), 11(3), 463-82 CODEN: CELLB5; ISSN: 0092-8674

(FILE 'HOME' ENTERED AT 13:49:09 ON 08 MAR 2005)

FILE 'STNGUIDE' ENTERED AT 13:49:13 ON 08 MAR 2005

FILE 'AGRICOLA, CAPLUS, BIOSIS' ENTERED AT 13:49:34 ON 08 MAR 2005 137 S SATELLITE TOBACCO MOSAIC VIRUS L10 S L1 AND ORIGIN OF ASSEMBLY L2L3 37 S L1 AND COAT PROTEIN L418 DUP REM L3 (19 DUPLICATES REMOVED) L5 316 S SATELLITE TOBACCO NECROSIS VIRUS L6 14 S L5 AND VECTOR L79 DUP REM L6 (5 DUPLICATES REMOVED) L8 46 S TOBACCO MOSAIC VIRUS AND ORIGIN OF ASSEMBLY 26 DUP REM L8 (20 DUPLICATES REMOVED) T.9

=> dup rem 110
PROCESSING COMPLETED FOR L10

#### => d 1-2 ti

- L11 ANSWER 1 OF 2 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- TI SVISS--a novel transient gene **silencing** system for gene function discovery and validation in tobacco plants.
- L11 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Recombinant expression system based on **satellite tobacco mosaic virus**

## -=> d ab

- L11 ANSWER 1 OF 2 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- We developed a novel, two-component transient gene silencing system in which the satellite tobacco mosaic virus (STMV) is used as vector for the delivery of inhibitory RNA into tobacco plants and the tobacco mosaic virus strain U2 (TMV-U2) is used as helper virus for supplying replication and movement proteins in trans. The main advantage of the system is that by uncoupling virus replication components from silencing induction components, the intensity of silencing becomes more pronounced. We call this system satellite virus-induced silencing system (SVISS) and will demonstrate here its robustness, speed and effectiveness. We were able to obtain pronounced and severe knockout phenotypes for a range of targeted endogenous genes belonging to various biochemical pathways and expressed in different plant tissues, such as genes involved in leaf and flower pigmentation, genes for cell wall synthesis in leaf, stem and root tissues or a ubiquitous RNA polymerase gene. By tandem insertion of more than one target gene sequence into the vector, we were able to induce simultaneous knockouts of an endogenous gene and a transgene. SVISS is the first transient gene silencing system for Nicotiana tabacum, which is a genetically well-characterized bridging species for the Solanaceae plant family.

#### => d so

- ANSWER 1 OF 2 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- SO The Plant journal: for cell and molecular biology, Dec 2002. Vol. 32, No. 5. p. 859-866
  Publisher: Oxford: Blackwell Sciences Ltd.
  ISSN: 0960-7412

.....

# => d 2 so

- L11 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN
- SO PCT Int. Appl., 48 pp.

CODEN: PIXXD2

# => d 2 pi

L11 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN PATENT NO. KIND DATE APPLICATION NO.

DATE

W: CA, JP, US RW: AT, BE, CH, DE, DK, ES, FR, GB, IT, LU, NL, SE

=> d 2 ab

ANSWER 2 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN

AB A cDNA encoding a full-length copy of the RNA genome of satellite

tobacco mosaic virus is cloned and

characterized for use as a transformation vector for plants. The virus is

systemic and needs a helper virus for replication and so can be used to

introduce foreign DNA into a plant without being pathogenic. The genome

was cloned as a cDNA by standard methods using defined oligonucleotide

primers. The RNA was shown by in vitro translation to encode two peptides

of mol. weight 6,800 and 17,500. A series of plasmid derivs. containing modified

cDNAs were constructed and transcripts from these used in infectivity

studies on tobacco plants with tobacco mosaic virus isolates as helpers.

The transcripts tested produced infected plants and virions were recovered

from these plants. Antisense transcripts were not infective.

=> d ti

- L12 ANSWER 1 OF 1 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN Different 5' leader sequences modulate beta-glucuronidase accumulation levels in transgenic Nicotiana tabacum plants.
- => s ((meulewaeter, f?) or (meulewaeter f?))/au L14 39 ((MEULEWAETER, F?) OR (MEULEWAETER F?))/AU

=> dup rem 115
PROCESSING COMPLETED FOR L15
L16 9 DUP REM L15 (9 DUPLICATES REMOVED)

=> d 1-9 ti

- L16 ANSWER 1 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- TI SVISS--a novel transient gene silencing system for gene function discovery and validation in tobacco plants.
- L16 ANSWER 2 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 2
  TI The 5' and 3' extremities of the satellite tobacco
  necrosis virus translational enhancer domain contribute
  differentially to stimulation of translation
- L16 ANSWER 3 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 3
  TI Functionality of the STNV translational enhancer domain correlates with affinity for two wheat germ factors
- L16 ANSWER 4 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 4
  TI Features of the autonomous function of the translational enhancer domain
  of satellite tobacco necrosis virus

- L16 ANSWER 5 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 5
- TI 5'- and 3'-sequences of satellite tobacco necrosis virus RNA promoting translation in tobacco.
- L16 ANSWER 6 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 6
- TI Expression of tobacco necrosis virus open reading frames 1 and 2 is sufficient for the replication of satellite necrosis virus.
- L16 ANSWER 7 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 7
- TI The 3' untranslated region of satellite tobacco necrosis virus RNA stimulates translation in vitro
- L16 ANSWER 8 OF 9 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Specificity of satellite activation by tobacco necrosis virus correlates with nucleic acid hybridization pattern between helper virus isolates
- L16 ANSWER 9 OF 9 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN TI GENOME STRUCTURE OF TOBACCO NECROSIS VIRUS STRAIN A.

# => d so

- L16 ANSWER 1 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- SO The Plant journal: for cell and molecular biology, Dec 2002. Vol. 32, No. 5. p. 859-866
  Publisher: Oxford: Blackwell Sciences Ltd.
  ISSN: 0960-7412

#### => d ab

- L16 ANSWER 1 OF 9 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- AB We developed a novel, two-component transient gene silencing system in which the satellite tobacco mosaic virus (STMV) is used as vector for the delivery of inhibitory RNA into tobacco plants and the tobacco mosaic virus strain U2 (TMV-U2) is used as helper virus for supplying replication and movement proteins in trans. The main advantage of the system is that by uncoupling virus replication components from silencing induction components, the intensity of silencing becomes more pronounced. We call this system satellite virus-induced silencing system (SVISS) and will demonstrate here its robustness, speed and effectiveness. We were able to obtain pronounced and severe knockout phenotypes for a range of targeted endogenous genes belonging to various biochemical pathways and expressed in different plant tissues, such as genes involved in leaf and flower pigmentation, genes for cell wall synthesis in leaf, stem and root tissues or a ubiquitous RNA polymerase gene. By tandem insertion of more than one target gene sequence into the vector, we were able to induce simultaneous knockouts of an endogenous gene and a transgene. SVISS is the first transient gene silencing system for Nicotiana tabacum, which is a genetically well-characterized bridging species for the Solanaceae plant family.

=> dup rem 118
PROCESSING COMPLETED FOR L18
L19 7 DUP REM L18 (10 DUPLICATES REMOVED)

=> d 1-7 ti

- L19 ANSWER 1 OF 7 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 1
- TI SVISS--a novel transient gene silencing system for gene function discovery and validation in tobacco plants.
- L19 ANSWER 2 OF 7 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 2
- TI The 5' and 3' extremities of the satellite tobacco necrosis virus translational enhancer domain contribute differentially to stimulation of translation
- L19 ANSWER 3 OF 7 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 3
- TI Functionality of the STNV translational enhancer domain correlates with affinity for two wheat germ factors
- L19 ANSWER 4 OF 7 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 4
- TI Features of the autonomous function of the translational enhancer domain of satellite tobacco necrosis virus
- L19 ANSWER 5 OF 7 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 5
- TI 5'- and 3'-sequences of satellite tobacco necrosis virus RNA promoting translation in tobacco.
- L19 ANSWER 6 OF 7 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 6
- TI Expression of tobacco necrosis virus open reading frames 1 and 2 is sufficient for the replication of satellite necrosis virus.
- L19 ANSWER 7 OF 7 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 7
- TI The 3' untranslated region of satellite tobacco necrosis virus RNA stimulates translation in vitro

=> d 5 ab

- L19 ANSWER 5 OF 7 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2005) on STN DUPLICATE 5
- AB The RNA of satellite tobacco necrosis
  virus (STNV) is a monocistronic messenger that lacks both a 5' cap
  and a 3' poly(A) tail. The STNV trailer contains an autonomous
  translational enhancer domain (TED) that promotes translation in vitro by
  more than one order of magnitude when combined with the 5'-terminal 173 nt
  of STNV RNA. We now show that the responsible sequence within the 5'
  region maps to the first 38 nt of the STNV RNA. Mutational analysis
  indicated that the primary sequence of the STNV 5' 38 nt and TED is
  important for translation stimulation in vitro, but did not reveal a role
  for the complementarity between the two. Translation of chimeric STNV-cat
  RNAs in tobacco protoplasts showed that TED promotes translation in vivo
  of RNAs lacking a cap and/or a poly(A) tail. Similar to in vitro,
  TED-dependent translation in tobacco was stimulated further by the STNV 5'

```
%
38 nt.
```

=> d 5 so

L19 ANSWER 5 OF 7 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

(2005) on STN DUPLICATE 5

SO The Plant journal : for cell and molecular biology, Apr 1998. Vol. 14, No. 2. p. 169-176

Publisher: Oxford : Blackwell Sciences Ltd.

ISSN: 0960-7412

=> s ((jacobs, j?) or (jacobs j))/au L20 3168 ((JACOBS, J?) OR (JACOBS J))/AU

=> s ((van eldik, g?) or (van eldik g?))/au L22 36 ((VAN ELDIK, G?) OR (VAN ELDIK G?))/AU

=> s 124 and (satellite tobacco mosaic virus or satellite tobacco necrosis virus)
L25 3 L24 AND (SATELLITE TOBACCO MOSAIC VIRUS OR SATELLITE TOBACCO
NECROSIS VIRUS)

=> dup rem 125
PROCESSING COMPLETED FOR L25
L26 1 DUP REM L25 (2 DUPLICATES REMOVED)

=> d ti

L26 ANSWER 1 OF 1 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

(2005) on STN DUPLICATE 1

TI SVISS--a novel transient gene silencing system for gene function discovery and validation in tobacco plants.